Inventor: BERNS et al.

Docket No.: PENDING00071861

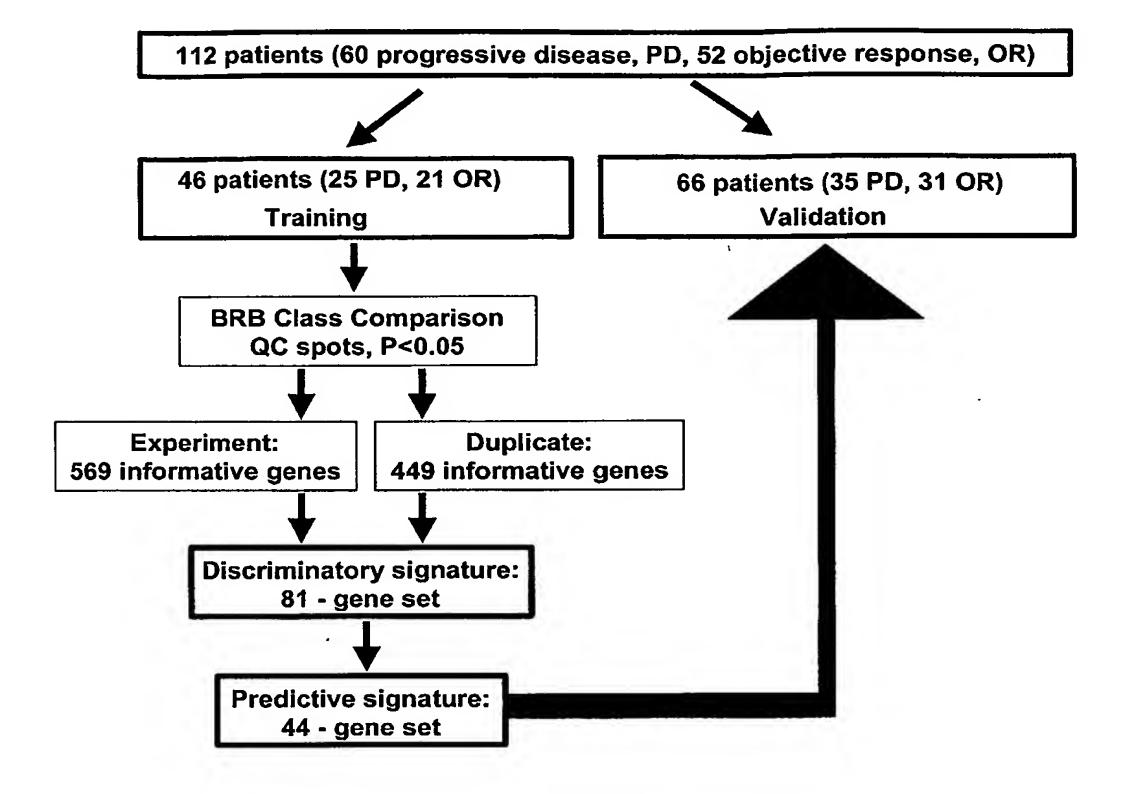
Title: PREDICTING RESPONSE AND OUTCOME OF METASTATIC BREAST CANCER

ANTI-ESTROGEN THERAPY

Serial No.: Unknown Sheet 1 of 5

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Figure 1



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Docket No.: PENDING00071861

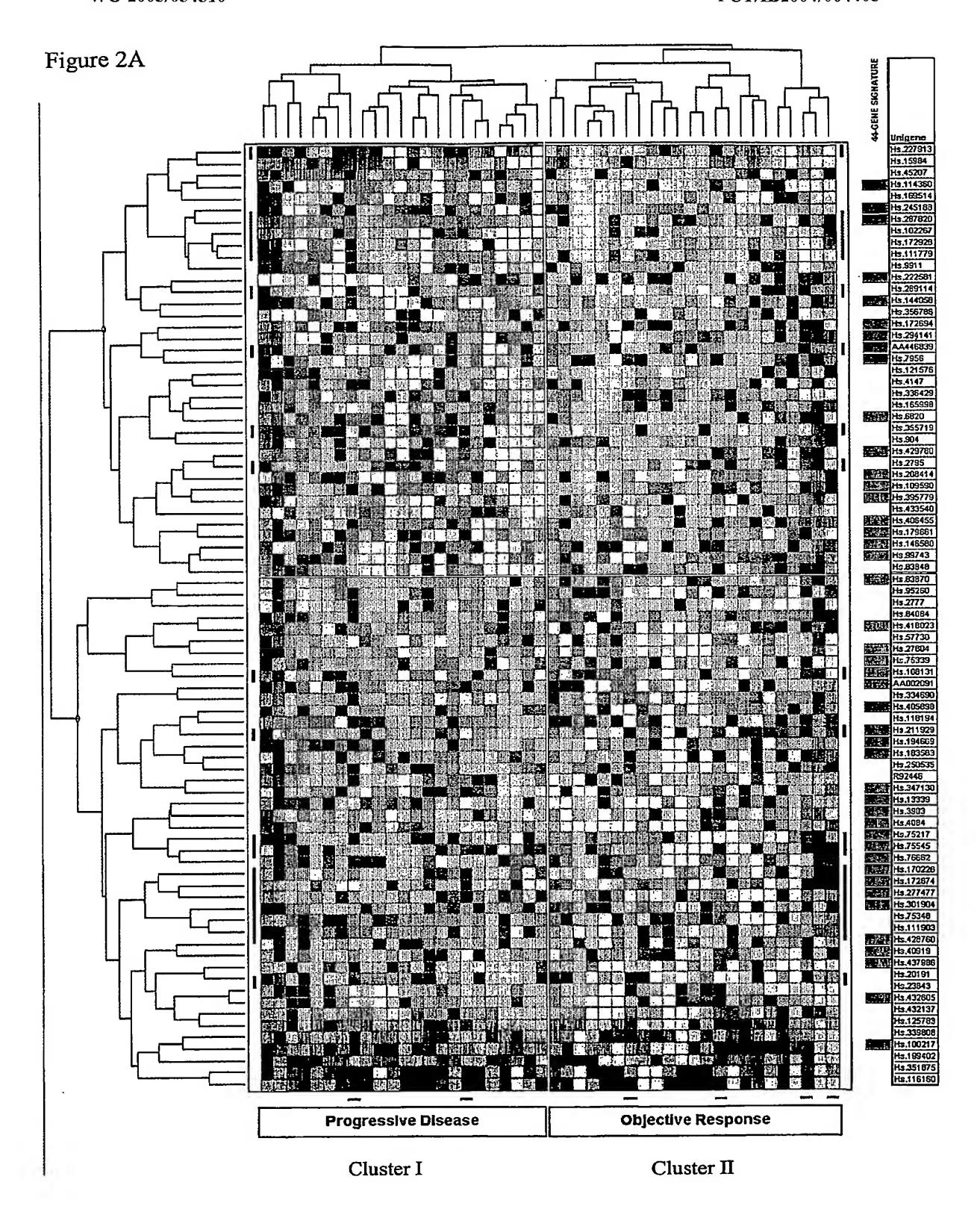
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ANTI-ESTROGEN THERAPY
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Figure 2B

Unigene Hs.227913 APIS 11pt2-q12 Hs.15994 LOCS1186 Xq22.1 Hs.16994 LOCS1186 Xq22.1 Hs.16914 Ks.245188 TMP3 22q12.3 Hs.169514 LOCS1186 Xq22.1 Hs.245188 TMP3 22q12.3 Metaloprotesse infator Hs.102267 LOX 523.2 Hs.192260 FM1 2q4 Cell adhesion Hs.102267 LOX 523.2 Extracellular matrix Hs.192260 FM1 2q4 Cell adhesion Hs.102267 LOX 523.2 Extracellular matrix Hs.111779 SPARC 5q31.3-q32 Extracellular matrix Hs.111779 SPARC 5q31.3-q32 Extracellular matrix Hs.14058 DKF2P564C103 17q45.2 Hs.144058 DKF2P564C103 17q45.2 Hs.151576 MYO18 2q12-q34 Myoshr, ATP-binding Actin-binding, Calmodulin-binding, Hs.165998 PAHRBP1 1p31-p32 Hs.155998 DABARAPL1 12p13.31 Endoplasmic relicular, Transmembrane, Olycopro, Hs.59319 PAHRBP1 1p31-p32 Hs.155998 CABARAPL1 12p13.31 Endoplasmic relicular, Transmembrane, Olycopro, Hs.5935 PAHRBP1 1p31-p32 Hs.165998 GEDX-3414 4q24-q25 Nuclear protein, Phosphorylation, RNA-binding, Ph.15691 15q21-q34 1p35-2 Cheperone. Hs.2795 LOHA 11p15.4 Oxidoreductase, NAD, Glycolysts Hs.19591 OKSW-cl.56 Gp21.33 Microtubuda, CTP-binding PAHRP 1p31-p32 Cheperone. Hs.19590 CNO2 12p13 Lyase; Olycolysts, Magnesiam	ER regulation 8 eln. 21 19
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Hs.15984 LOCS1185 Xq22.1	ein. 21
Hs. 45207 CHD6-pending 20q12 Transcription regulation; Repressor; Nuclear proteins Hs. 14350 TSC22 13q14 Transcription regulation; Repressor; Nuclear proteins Hs. 245188 TMP3 22q12.3 Metalloprotesse inhibitor. Cell edhesion. Hs. 102267 LOX 5q23.2 Extracellular matrix. Hs. 111779 SPARC 5q31.3-q32 Extracellular matrix. Hs. 111779 SPARC 5q31.3-q32 Extracellular matrix. Hs. 111779 SPARC 5q31.3-q32 Extracellular matrix. Hs. 19311 TNC 5q33 Extracellular matrix. Hs. 144058 DKF2P564C103 17q25.2 Hs. 359788 GNAQ 9q21.2 Hs. 172694 Hs. 172694 LOC117584 17q12 Hs. 172694 Hs. 174059 LOC117584 17q12 Hs. 172694 Hs. 172695 Hs. 172	ein. 21
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Hs.289114 TNC	
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	id biosynthesis.
Hs.83970 NEB 2q22 Actin-binding Muscle protein Cytoskeleton.	
Hs.95260 FAM6A1 6p22-p23	
Hs.2777 ITUH1 3p21.2-p21.1 Serine protesse inhibitor. Hs.84084 APPBP2 17q21-q23	
Hs.418023	
Hs.57730 KIAA0133 1q42.13 Hypothetical protein; Transmembrane.	
Hs.27804	
Hs.75339 INPPL1 11q23	
Hs.108131 CASP2 7q34-q35 Hydrolese; Thiol proteese; Apoptosis.	
AAU02091 CACH-1 5q14.1	
Hs.334690	
Hs.405898 KIAA0999 11q23.3	
Hs.118194 DBR1 3q22.3	ndrion. 8
Hs.211929 TXN2 22q13.1 Redox-active center, Electron transport, M3ochor Ball Hs.194669 EZH1 17q21.1-q21.3 Transcription regulation, Nuclear protein, DNA-bin	
Hs.194669 EZH1 17q21.1-q21.3 Transcription regulation, Nuclear protein, DNA-bin Hs.183583 SERPINE1 6p25 Serpin, Serine protease Inhibitor.	
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R92448	
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Hs.13339 PRPSAP2 17p11.2-p12	
Hs.3903 CDC42EP4 17q24-q25 Cytoskeleton.	21
程 Hs.4084 KIAA1025 12q24.22	
Hs.75217 MAP2K4 17p11.2 Serine/liveonins-protein kinese; Tyrosins-protein	китаза.
Hs.75545 E.4R 16p11.2-12.1 Receptor; Transmembrane; Glycoprotein.	
Hs.76662 APH2 10q24.1 Hs.170226 FL38045 9q33.1	
Hs.17026 FL/38045 9q33.1 Hs.172674 NFATC3 16q22.2 Transcription regulation; Activator; Nuclear protein	n: DNA-binding.
His 277477 HLA-C 6p21.3 MHC t, Transmembrane; Glycoprotein.	18
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Hs.40918 FLJ14511 9q22.33	
Ha.437986	<u></u>
Hs.20191 SIAH2 3q25	8
Hs.23643 MST4 Xq28.1 Hs.432605 UGCO 8q31 Glycosyltrensferese; Endoplasmic reticulum.	17
Hs.432605 UGCO 9q31 Gtycosyttransferese; Endoplasmic reticulum. Hs.432137 DUX2	
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Hs.339808 KIAA0563 17q21.31	
Hs.100217 FMNL 17q21	6
Hs.199402 UFD1L 22q11.21 Ubl conjugation pethway.	- V
Hs.351875 COX6C 8q22-q23 Oxidoreductase; Inner membrane; Mitochondrion.	
Hs.116160	

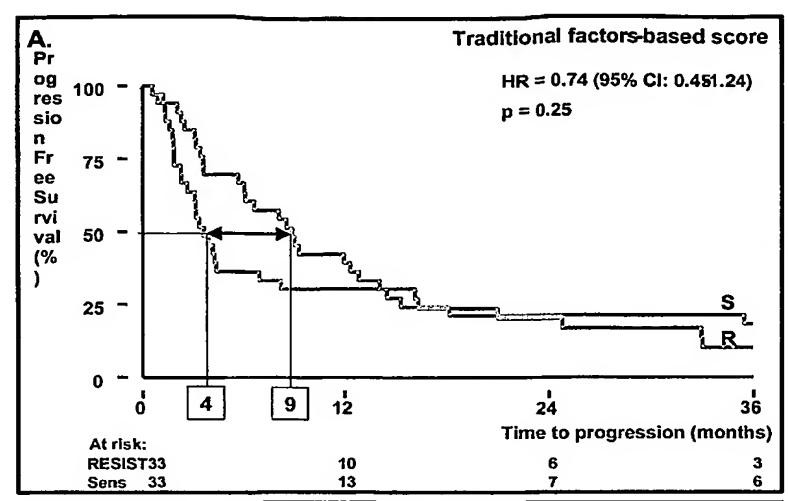
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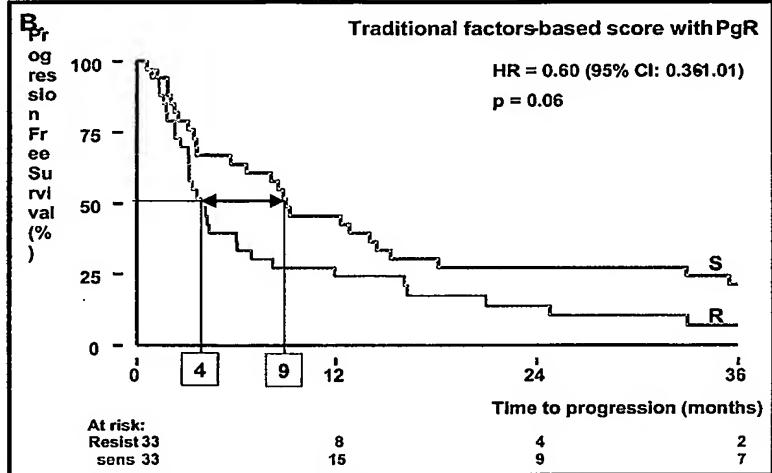
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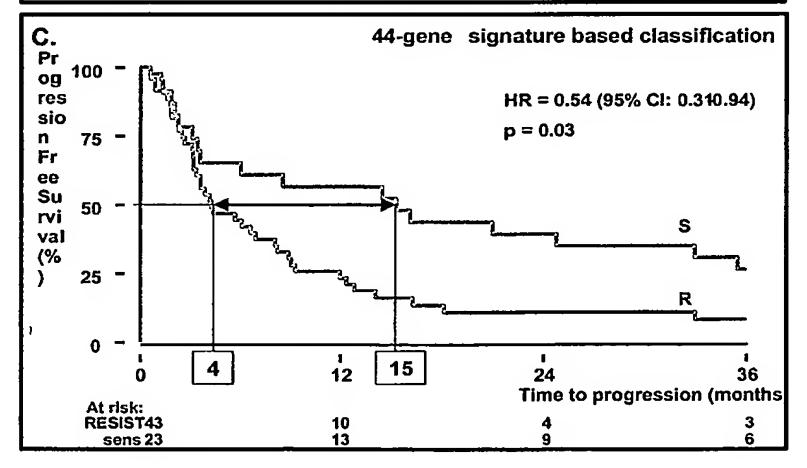
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Figure 3







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